

SEQUENCE LISTING

<110> Applied Reasearch Systems ARS holding

<120> NOVEL OX40R BINDING AGENTS

<130> W0498

<160> 15

<170> PatentIn version 3.0

<210> 1

<211> 183

<212> PRT

<213> Homo sapiens

<400> 1

```

Met Glu Arg Val Gln Pro Leu Glu Glu Asn Val Gly Asn Ala Ala Arg
1      5      10
Pro Arg Phe Glu Arg Asn Lys Leu Leu Leu Val Ala Ser Val Ile Gln
20     25     30
Gly Leu Gly Leu Leu Leu Cys Phe Thr Tyr Ile Cys Leu His Phe Ser
35     40     45
Ala Leu Gln Val Ser His Arg Tyr Pro Arg Ile Gln Ser Ile Lys Val
50     55     60
Gln Phe Thr Glu Tyr Lys Lys Glu Lys Gly Phe Ile Leu Thr Ser Gln
65     70     75     80
Lys Glu Asp Glu Ile Met Lys Val Gln Asn Asn Ser Val Ile Ile Asn
85     90     95
Cys Asp Gly Phe Tyr Leu Ile Ser Leu Lys Gly Tyr Phe Ser Gln Glu
100    105    110
Val Asn Ile Ser Leu His Tyr Gln Lys Asp Glu Glu Pro Leu Phe Gln
115    120    125
Leu Lys Lys Val Arg Ser Val Asn Ser Leu Met Val Ala Ser Leu Thr
130    135    140
Tyr Lys Asp Lys Val Tyr Leu Asn Val Thr Thr Asp Asn Thr Ser Leu
145    150    155    160
Asp Asp Phe His Val Asn Gly Gly Glu Leu Ile Leu Ile His Gln Asn
165    170    175

```

Pro Gly Glu Phe Cys Val Leu
180

<210> 2

<211> 31

<212> PRT

<213> synthetic construct

<400> 2

Val Ala Ser Leu Thr Tyr Lys Asp Lys Val Tyr Leu Asn Val Thr Thr
1 5 10 15

Asp Asn Thr Ser Leu Asp Asp Phe His Val Asn Gly Gly Glu Leu
20 25 30

<210> 3

<211> 24

<212> PRT

<213> synthetic construct

<400> 3

Leu Asp Asp Phe His Val Asn Gly Gly Glu Leu Ile Leu Ile His Gln
1 5 10 15

Asn Pro Gly Glu Phe Cys Val Leu
20

<210> 4

<211> 29

<212> PRT

<213> synthetic construct

<400> 4

Val Ser His Arg Tyr Pro Arg Ile Gln Ser Ile Lys Val Gln Phe Thr
1 5 10 15

Glu Tyr Lys Lys Glu Lys Gly Phe Ile Leu Thr Ser Gln
20 25

<210> 5

<211> 31

<212> PRT

<213> synthetic construct

<400> 5

Glu Lys Gly Phe Ile Leu Thr Ser Gln Lys Glu Asp Glu Ile Met Lys
 1 5 10 15

Val Gln Asn Asn Ser Val Ile Ile Asn Cys Asp Gly Phe Tyr Leu
 20 25 30

<210> 6

<211> 31

<212> PRT

<213> synthetic construct

<400> 6

Ile Ile Asn Cys Asp Gly Phe Tyr Leu Ile Ser Leu Lys Gly Tyr Phe
 1 5 10 15

Ser Gln Glu Val Asn Ile Ser Leu His Tyr Gln Lys Asp Glu Glu
 20 25 30

<210> 7

<211> 30

<212> PRT

<213> synthetic construct

<400> 7

His Tyr Gln Lys Asp Glu Glu Pro Leu Phe Gln Leu Lys Lys Arg Ser
 1 5 10 15

Val Asn Ser Leu Met Val Ala Ser Leu Thr Tyr Lys Asp Lys
 20 25 30

<210> 8

<211> 10

<212> PRT

<213> synthetic construct

<400> 8

Gly Tyr Phe Ser Gln Glu Val Asn Ile Ser
 1 5 10

<210> 9

<211> 10

<212> PRT

<213> synthetic construct

<400> 9

| | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Ile | Ser | Leu | His | Tyr | Gln | Lys | Asp | Glu | Glu |
| 1 | | | | 5 | | | | | 10 |

<210> 10

<211> 10

<212> PRT

<213> synthetic construct

<400> 10

| | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Gly | Phe | Tyr | Leu | Ile | Ser | Leu | Lys | Gly | Tyr |
| 1 | | | | 5 | | | | | 10 |

<210> 11

<211> 10

<212> PRT

<213> synthetic construct

<400> 11

| | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Gln | Glu | Val | Asn | Ile | Ser | Leu | His | Tyr | Gln |
| 1 | | | | 5 | | | | | 10 |

<210> 12

<211> 10

<212> PRT

<213> synthetic construct

<400> 12

| | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Ile | Ile | Asn | Cys | Asp | Gly | Phe | Tyr | Leu | Ile |
| 1 | | | | 5 | | | | | 10 |

<210> 13

<211> 5

<212> PRT

<213> synthetic construct

<400> 13

| | | | | |
|-----|-----|-----|-----|-----|
| Gly | Tyr | Phe | Ser | Gln |
| 1 | | | | 5 |

<210> 14

<211> 18

<212> PRT

<213> synthetic construct

<400> 14

Leu Lys Gly Ser Phe Phe Gln Glu Val Lys Ile Asp Leu His Phe Arg
1 5 10 15

Glu Asp
20

<210> 15

<211> 18

<212> PRT

<213> synthetic construct

<400> 15

Ala Phe Lys Asp Lys Val Tyr Leu Thr Val Asn Ala Pro Asp Thr Leu
1 5 10 15

Cys Glu
20